

PCT

RAW SEQUENCE LISTING DATE: 10/08/2004 PATENT APPLICATION: US/10/509,975 TIME: 13:57:37

Input Set : A:\2543-1-036PCTUS - Seq listing.txt

Output Set: N:\CRF4\10082004\J509975.raw

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3 <110> APPLICANT: Patel, Sonal
      5 <120> TITLE OF INVENTION: SC6 FOR DIAGNOSIS OF HYPOXIA RELATED CONDITIONS
      7 <130> FILE REFERENCE: 2543-1-036PCT/US
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/509,975
                                                                  THE CONTROL OF THE STATE OF THE STATE OF
C--> 10 <141> CURRENT FILING DATE: 2004-10-01
     12 <150> PRIOR APPLICATION NUMBER: GB 0207533.1
     13 <151> PRIOR FILING DATE: 2002-04-02
     15 <160> NUMBER OF SEQ ID NOS: 4
     17 <170> SOFTWARE: PatentIn version 3.1
     19 <210> SEQ ID NO: 1
     20 <211> LENGTH: 619
     21 <212> TYPE: PRT
     22 <213> ORGANISM: Homo Sapiens
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     34 Glu Ala Glu Gly Lys Pro Pro Gln Arg Glu Lys Trp Ser Ser Lys Ile
     38 Asp Phe Val Leu Ser Val Ala Gly Gly Phe Val Gly Leu Gly Asn Val
     42 Trp Arg Phe Pro Tyr Leu Cys Tyr Lys Asn Gly Gly Gly Ala Phe Leu
                            70
     46 Ile Pro Tyr Phe Ile Phe Leu Phe Gly Ser Gly Leu Pro Val Phe Phe
     50 Leu Glu Ile Ile Ile Gly Gln Tyr Thr Ser Glu Gly Gly Ile Thr Cys
                    100
                                         105
     54 Trp Glu Lys Ile Cys Pro Leu Phe Ser Gly Ile Gly Tyr Ala Ser Val
                                     120
                                                         125
                115
     58 Val Ile Val Ser Leu Leu Asn Val Tyr Tyr Ile Val Ile Leu Ala Trp
                                135
     62 Ala Thr Tyr Tyr Leu Phe Gln Ser Phe Gln Lys Glu Leu Pro Trp Ala
                            150
                                                                      160
     66 His Cys Asn His Ser Trp Asn Thr Pro His Cys Met Glu Asp Thr Met
                                             170
     70 Arg Lys Asn Lys Ser Val Trp Ile Thr Ile Ser Ser Thr Asn Phe Thr
                                         185
                    180
     74 Ser Pro Val Ile Glu Phe Trp Glu Arq Asn Val Leu Ser Leu Ser Pro
                                     200
     78 Gly Ile Asp His Pro Gly Ser Leu Lys Trp Asp Leu Ala Leu Cys Leu
                                215
```

82 Leu Leu Val Trp Leu Val Cys Phe Phe Cys Ile Cys Lys Gly Val Arg

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	225					230					235					240
86	Ser	Thr	Gly	Lys	Val	Val	Tyr	Phe	Thr	Ala	Thr	Phe	Pro	Phe	Ala	Met
87					245					250					255	
90	Leu	Leu	Val	Leu	Leu	Val	Arg	Gly	Leu	Thr	Leu	Pro	Gly	Ala	Gly	Arg
91				260					265					270		
94	Glv	Ile	Lys	Phe	Tyr	Leu	Tyr	Pro	Asp	Ile	Thr	Arg	Leu	Glu	Asp	Pro
95	-		275		•		•	280				_	285		_	
	Gln	Val	Trp	Ile	Asp	Ala	Glv	Thr	Gln	Ile	Phe	Phe	Ser	Tvr	Ala	Ile
99		290					295					300		-1-		
) (Tyre		1 G13	, Ala	Met	Thr		^ T.e.ı	ı Gly	z Sei	ר דעז		Lvs	: ጥህን	Tive	Tyr
	305		. 01	HIG	· NC	310			. 01	, 00.	315		. <u>.</u> .,.	, -,-		320
			c Tun	^ Arc	λer			· T.e1	1 T.A1	, G1s			ı Acr	Ser	- G1v	Thr
		1 261	. тул	. ALG	325		, MC	. пес	и шес	330		, Let	LASI	1 561	335	
107		Dha	. 37-1	60*			- דת	. т1-	Dhe			La	. (2)	, Dhe		Ala
		PILE	e val		_	PILE	: Alc	, TTG			. 116	: шес	r GT			. AIA
111		. al.		340			. ~1.		345					350		
		1 GIL			val	LASE	TIE			ya.	L Alč	i GI			PIC	Gly
115			355				_	360					365			
				e lle	AL	а Туг			S Ala	a va.	ı Tnı			: Pro	о тел	Pro
119		370					375				_	380		_		_
122	? Thr	Phe	e Trr	Ser	Ile			e Phe	e Ile	e Met			ı Leı	ı Let	ı GIŞ	Leu
	385					390				_	395				_	400
126	Asp	Sei	c Glr	ı Phe	· Val	l Gli	ı Val	. Gli	ı Gly	/ Gli	n Ile	e Thi	Sei	: Let	ı Val	. Asp
127					405					41(415	
130) Let	туз	r Pro	Ser	Phe	e Lei	ı Arg	J Lys	s Gly	у Туі	r Arg	g Arg	g Gli	ı Ile	e Ph∈	lle
131	_			420					425	5 .				430)	
134	Ala	Phe	e Val	L Cys	Sei	: Ile	e Sei	туз	r Lei	ı Leı	ı Gly	/ Let	ı Thi	: Met	. Val	Thr
135	5 ·		435	5				44()				445	5		
138	Glu	Gly	y Gly	/ Met	Туз	r Val	l Phe	Glr	ı Leı	ı Phe	e Asp	ту Ту	туз:	: Ala	a Ala	. Ser
139)	450)				455	5				460)			
142	Gly	va.	l Cys	. Leu	. Le	ı Trp	Va]	Ala	a Phe	e Phe	e Glu	ı Cys	Phe	e Val	l Ile	. Ala
143	465	;				470)				475	5				480
146	Tr	Ile	e Tyr	Gly	Gly	Asp	Asr	ı Leı	י Tyı	r Ası	Gly	/ Ile	e Glu	ı Asp	Met	: Ile
147	_		-	_	485					490					495	
150) Gly	Ty:	r Arc	, Pro	Gly	/ Pro	Tr	Met	Ly:	з Туг	c Sei	rrr	Va]	Ile	Thr	Pro
151	-	-	•	500	_	•	•		50					510		
		. Leı	ı Cvs	. Val	Glv	/ Cvs	s Phe	e Ile	e Phe	e Sei	r Lei	ı Val	Lys	Ty:	. Val	Pro
155			515		- 4			520					525			
		Thi			Lvs	Thr	TVI			r Pro	Thi	rr	Ala	a Ile	e Gly	Leu
159		530	-		1		535					540				
				r T.e.	ıΑla	a T.e.			r Met	. Lei	1 Cvs			Lei	ı Val	. Ile
	545		, ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,			550					555					560
			- Arc	T.A.1	. (376			^ Gl1	1 G11	, Dro			ı Val	Arc	r Val	. Lys
167			- 27.7	, 1100	565			. 51	. 01)	570			_ , , , ,		575	
		. T.o.	. T.	. The			· Cl.	Dro	. A.c.			. 77-	. Wal	G1,		
	_	. пе	י הפו	580		, WIG	9 91	. FI	585		3 TTF	, WIG	, va.	590		g Glu
171		. אT.	, mb-			~ 7 ~-		- 71 m-			Ma+	- 7 ~~	, cı.			Wal
	_	. WT			, туг	. Abi	. 561	600		L val	ואפנ	. ASI	609		י חבו	ı Val
175		. D	595			. 77-	, TT-7			~ N/~ 1		_	80:	,		
	-			r His	; TT6	: 116			ı III	т мет	. Met	-				
179	,	610	J				615	•								

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185 <213> ORGANISM: Homo Sapiens
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190 caaggacatg gtgaagcct caccagggaa gagcccaggc acgcggcctg aggacgaggc
                                                                        180
192 tgagggaaaa cctccgcaga gggagaagtg gtctagcaag atcgactttg tgctctctgt
194 ggctggcggc ttcgtgggct tgggcaacgt ctggcgcttc ccgtacctct gctacaagaa
                                                                        240
196 tggtggaggt gcgtttctca taccgtattt tattttcctg tttgggagcg gcctgcctgt.
                                                                        300
198 gtttttcttg gagatcatca taggccagta cacctctgaa gggggcatca cctgctggga
                                                                        360
200 aaagatetge ceettgttet etggtategg etatgeetee gttgtaattg tgteeeteet
                                                                        420
202 qaatqtctac tacatcqtca tcctgqcctg ggccacatac tacctgttcc agtccttcca
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206 caccatgogo aagaacaaga gtgtctggat caccatcago tocaccaact tcacctcocc
                                                                        600
208 tgtcatcgag ttctgggagc gcaacgtgct gagcttgtcc cctggaatcg accacccagg
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210 ctctctgaaa tgggacctcg ctctctgcct tcttttagtc tggctagtgt gtttcttctg
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212 catctgcaag ggcgtcaggt ccactgggaa ggtcgtctac ttcacagcca cttttccatt
                                                                        780
214 egecatgete etggtgetge tggteegagg getgaegetg eegggegegg geegaggeat
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216 caagttotat ctgtatcctg acatcacccg ccttgaggac ccacaggtgt ggattgacgc
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218 tgggactcag atattettet ettatgeeat etgeetgggg getatgacet egetggggag
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220 ctacaacaag tacaagtata actcgtacag ggactgtatg ctgctgggat gcctgaacag
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1140
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232 cgtgtgtagc atcagctacc tgctggggct gacgatggtg acggagggtg gcatgtatgt
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264 tcattttcaa aagcaattct tcggtgctgt gtagctggca gaaagttctg tccagtaaac
266 gcaggatgga attttcctgg gactctacac ccatcttaag gtggtatacc ttccaaatcc
                                                                       2400
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268 tggttcagat ggaagaaata gcaggagaga ggacccatta gctggcagac ccaggggaag
                                                                       2520
270 aaaggagggc tgtgaggaga tacctcatta aacttggctt agtgaagaag agagatgcca
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272 aaggaatgaa ccaaccette acataaagga gactggctga agetgaatga ggaggceeta
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276	aattttgaga tggtga	gtgg	atagtcagta	gaccgtcaga	accactggcc	agagaggag	2700			
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294	cctccactgg ggaagt	gctc	ccagttcaga	acaagggcag	cccgtggtgc	tgacctagga	3240			
296	tataacaaag ctcttc	actt	caaaacccct	gcaatagctg	ggtttacaga	catttaccac	3300			
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VERIFICATION SUMMARY

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L:9 M:270 C: Current Application Number differs, Replaced Current Application Number

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date